# RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/582,973A_

Source: _1FWO_

Date Processed by STIC: _12/5/06_

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RAW SEQUENCE LISTING                          DATE: 12/05/2006
PATENT APPLICATION:  US/10/582,973A          TIME: 09:48:41

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   3 <110> APPLICANT: NAKAMURA, Toshikazu
   4         MATSUMOTO, Kunio
   5         FUKUTA, Kazuhiro
   7 <120> TITLE OF INVENTION: GLYCOSYLATION-DEFICIENT HEPATOCYTE GROWTH FACTOR
   9 <130> FILE REFERENCE: 2006_0825A
  11 <140> CURRENT APPLICATION NUMBER: 10/582,973A
C-->  12 <141> CURRENT FILING DATE: 2006-06-15
  14 <150> PRIOR APPLICATION NUMBER: PCT/JP04/18719
  15 <151> PRIOR FILING DATE: 2004-12-15
  17 <150> PRIOR APPLICATION NUMBER: JP 2003-418790
  18 <151> PRIOR FILING DATE: 2003-12-16
  20 <150> PRIOR APPLICATION NUMBER: JP 2003-425691
  21 <151> PRIOR FILING DATE: 2003-12-22
  23 <160> NUMBER OF SEQ ID NOS: 8
  25 <170> SOFTWARE: PatentIn version 3.3
  27 <210> SEQ ID NO: 1
  28 <211> LENGTH: 728
  29 <212> TYPE: PRT
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  43         35                  40                  45
  46 Thr Leu Ile Lys Ile Asp Pro Ala Leu Lys Ile Lys Thr Lys Lys Val
  47     50                  55                  60
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  51 65                  70                  75                  80
  54 Pro Phe Thr Cys Lys Ala Phe Val Phe Asp Lys Ala Arg Lys Gln Cys
  55             85                  90                  95
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  59             100                 105                 110
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  63         115                 120                 125
  66 Ile Ile Gly Lys Gly Arg Ser Tyr Lys Gly Thr Val Ser Ile Thr Lys
  67     130                 135                 140
  70 Ser Gly Ile Lys Cys Gln Pro Trp Ser Ser Met Ile Pro His Glu His
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 83         195               200               205
 86 Val Glu Cys Met Thr Cys Asn Gly Glu Ser Tyr Arg Gly Leu Met Asp
 87     210               215               220
 90 His Thr Glu Ser Gly Lys Ile Cys Gln Arg Trp Asp His Gln Thr Pro
 91 225               230               235               240
 94 His Arg His Lys Phe Leu Pro Glu Arg Tyr Pro Asp Lys Gly Phe Asp
 95               245               250               255
 98 Asp Asn Tyr Cys Arg Asn Pro Asp Gly Gln Pro Arg Pro Trp Cys Tyr
 99               260               265               270
102 Thr Leu Asp Pro His Thr Arg Trp Glu Tyr Cys Ala Ile Lys Thr Cys
103         275               280               285
106 Ala Asp Asn Thr Met Asn Asp Thr Asp Val Pro Leu Glu Thr Thr Glu
107     290               295               300
110 Cys Ile Gln Gly Gln Gly Glu Gly Tyr Arg Gly Thr Val Asn Thr Ile
111 305               310               315               320
114 Trp Asn Gly Ile Pro Cys Gln Arg Trp Asp Ser Gln Tyr Pro His Glu
115               325               330               335
118 His Asp Met Thr Pro Glu Asn Phe Lys Cys Lys Asp Leu Arg Glu Asn
119         340               345               350
122 Tyr Cys Arg Asn Pro Asp Gly Ser Glu Ser Pro Trp Cys Phe Thr Thr
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126 Asp Pro Asn Ile Arg Val Gly Tyr Cys Ser Gln Ile Pro Asn Cys Asp
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131 385               390               395               400
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135               405               410               415
138 Lys Asn Met Glu Asp Leu His Arg His Ile Phe Trp Glu Pro Asp Ala
139         420               425               430
142 Ser Lys Leu Asn Glu Asn Tyr Cys Arg Asn Pro Asp Asp Asp Ala His
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146 Gly Pro Trp Cys Tyr Thr Gly Asn Pro Leu Ile Pro Trp Asp Tyr Cys
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150 Pro Ile Ser Arg Cys Glu Gly Asp Thr Thr Pro Thr Ile Val Asn Leu
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154 Asp His Pro Val Ile Ser Cys Ala Lys Thr Lys Gln Leu Arg Val Val
155               485               490               495
158 Asn Gly Ile Pro Thr Arg Thr Asn Ile Gly Trp Met Val Ser Leu Arg
159         500               505               510
162 Tyr Arg Asn Lys His Ile Cys Gly Gly Ser Leu Ile Lys Glu Ser Trp
163         515               520               525
166 Val Leu Thr Ala Arg Gln Cys Phe Pro Ser Arg Asp Leu Lys Asp Tyr
167     530               535               540
170 Glu Ala Trp Leu Gly Ile His Asp Val His Gly Arg Gly Asp Glu Lys
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182 Phe Val Ser Thr Ile Asp Leu Pro Asn Tyr Gly Cys Thr Ile Pro Glu
183         595             600             605
186 Lys Thr Ser Cys Ser Val Tyr Gly Trp Gly Tyr Thr Gly Leu Ile Asn
187     610             615             620
190 Tyr Asp Gly Leu Leu Arg Val Ala His Leu Tyr Ile Met Gly Asn Glu
191 625             630             635             640
194 Lys Cys Ser Gln His His Arg Gly Lys Val Thr Leu Asn Glu Ser Glu
195             645             650             655
198 Ile Cys Ala Gly Ala Glu Lys Ile Gly Ser Gly Pro Cys Glu Gly Asp
199         660             665             670
202 Tyr Gly Gly Pro Leu Val Cys Glu Gln His Lys Met Arg Met Val Leu
203         675             680             685
206 Gly Val Ile Val Pro Gly Arg Gly Cys Ala Ile Pro Asn Arg Pro Gly
207     690             695             700
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219 <211> LENGTH: 723
220 <212> TYPE: PRT
221 <213> ORGANISM: Homo sapiens
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229 Leu His Leu Leu Leu Leu Pro Ile Ala Ile Pro Tyr Ala Glu Gly Gln
230         20              25              30
233 Arg Lys Arg Arg Asn Thr Ile His Glu Phe Lys Lys Ser Ala Lys Thr
234         35              40              45
237 Thr Leu Ile Lys Ile Asp Pro Ala Leu Lys Ile Lys Thr Lys Lys Val
238     50              55              60
241 Asn Thr Ala Asp Gln Cys Ala Asn Arg Cys Thr Arg Asn Lys Gly Leu
242 65              70              75              80
245 Pro Phe Thr Cys Lys Ala Phe Val Phe Asp Lys Ala Arg Lys Gln Cys
246             85              90              95
249 Leu Trp Phe Pro Phe Asn Ser Met Ser Ser Gly Val Lys Lys Glu Phe
250         100             105             110
253 Gly His Glu Phe Asp Leu Tyr Glu Asn Lys Asp Tyr Ile Arg Asn Cys
254     115             120             125
257 Ile Ile Gly Lys Gly Arg Ser Tyr Lys Gly Thr Val Ser Ile Thr Lys
258     130             135             140
261 Ser Gly Ile Lys Cys Gln Pro Trp Ser Ser Met Ile Pro His Glu His
262 145             150             155             160
265 Ser Tyr Arg Gly Lys Asp Leu Gln Glu Asn Tyr Cys Arg Asn Pro Arg
266             165             170             175
269 Gly Glu Glu Gly Gly Pro Trp Cys Phe Thr Ser Asn Pro Glu Val Arg
270         180             185             190
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273 Tyr Glu Val Cys Asp Ile Pro Gln Cys Ser Glu Val Glu Cys Met Thr
274         195                 200                 205
277 Cys Asn Gly Glu Ser Tyr Arg Gly Leu Met Asp His Thr Glu Ser Gly
278         210                 215                 220
281 Lys Ile Cys Gln Arg Trp Asp His Gln Thr Pro His Arg His Lys Phe
282 225                 230                 235                 240
285 Leu Pro Glu Arg Tyr Pro Asp Lys Gly Phe Asp Asp Asn Tyr Cys Arg
286             245                 250                 255
289 Asn Pro Asp Gly Gln Pro Arg Pro Trp Cys Tyr Thr Leu Asp Pro His
290             260                 265                 270
293 Thr Arg Trp Glu Tyr Cys Ala Ile Lys Thr Cys Ala Asp Asn Thr Met
294         275                 280                 285
297 Asn Asp Thr Asp Val Pro Leu Glu Thr Thr Glu Cys Ile Gln Gly Gln
298         290                 295                 300
301 Gly Glu Gly Tyr Arg Gly Thr Val Asn Thr Ile Trp Asn Gly Ile Pro
302 305                 310                 315                 320
305 Cys Gln Arg Trp Asp Ser Gln Tyr Pro His Glu His Asp Met Thr Pro
306             325                 330                 335
309 Glu Asn Phe Lys Cys Lys Asp Leu Arg Glu Asn Tyr Cys Arg Asn Pro
310             340                 345                 350
313 Asp Gly Ser Glu Ser Pro Trp Cys Phe Thr Thr Asp Pro Asn Ile Arg
314         355                 360                 365
317 Val Gly Tyr Cys Ser Gln Ile Pro Asn Cys Asp Met Ser His Gly Gln
318         370                 375                 380
321 Asp Cys Tyr Arg Gly Asn Gly Lys Asn Tyr Met Gly Asn Leu Ser Gln
322 385                 390                 395                 400
325 Thr Arg Ser Gly Leu Thr Cys Ser Met Trp Asp Lys Asn Met Glu Asp
326             405                 410                 415
329 Leu His Arg His Ile Phe Trp Glu Pro Asp Ala Ser Lys Leu Asn Glu
330             420                 425                 430
333 Asn Tyr Cys Arg Asn Pro Asp Asp Asp Ala His Gly Pro Trp Cys Tyr
334         435                 440                 445
337 Thr Gly Asn Pro Leu Ile Pro Trp Asp Tyr Cys Pro Ile Ser Arg Cys
338         450                 455                 460
341 Glu Gly Asp Thr Thr Pro Thr Ile Val Asn Leu Asp His Pro Val Ile
342 465                 470                 475                 480
345 Ser Cys Ala Lys Thr Lys Gln Leu Arg Val Val Asn Gly Ile Pro Thr
346             485                 490                 495
349 Arg Thr Asn Ile Gly Trp Met Val Ser Leu Arg Tyr Arg Asn Lys His
350             500                 505                 510
353 Ile Cys Gly Gly Ser Leu Ile Lys Glu Ser Trp Val Leu Thr Ala Arg
354         515                 520                 525
357 Gln Cys Phe Pro Ser Arg Asp Leu Lys Asp Tyr Glu Ala Trp Leu Gly
358         530                 535                 540
361 Ile His Asp Val His Gly Arg Gly Asp Glu Lys Cys Lys Gln Val Leu
362 545                 550                 555                 560
365 Asn Val Ser Gln Leu Val Tyr Gly Pro Glu Gly Ser Asp Leu Val Leu
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369 Met Lys Leu Ala Arg Pro Ala Val Leu Asp Asp Phe Val Ser Thr Ile
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370          580              585              590
373 Asp Leu Pro Asn Tyr Gly Cys Thr Ile Pro Glu Lys Thr Ser Cys Ser
374          595              600              605
377 Val Tyr Gly Trp Gly Tyr Thr Gly Leu Ile Asn Tyr Asp Gly Leu Leu
378      610              615              620
381 Arg Val Ala His Leu Tyr Ile Met Gly Asn Glu Lys Cys Ser Gln His
382 625              630              635              640
385 His Arg Gly Lys Val Thr Leu Asn Glu Ser Glu Ile Cys Ala Gly Ala
386              645              650              655
389 Glu Lys Ile Gly Ser Gly Pro Cys Glu Gly Asp Tyr Gly Gly Pro Leu
390          660              665              670
393 Val Cys Glu Gln His Lys Met Arg Met Val Leu Gly Val Ile Val Pro
394          675              680              685
397 Gly Arg Gly Cys Ala Ile Pro Asn Arg Pro Gly Ile Phe Val Arg Val
398      690              695              700
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419 gaattcaaaa aatcagcaaa gactacccta atcaaaatag atccagcact gaagataaaa      180
421 accaaaaaag tgaatactgc agaccaatgt gctaatagat gtactaggaa taaaggactt      240
423 ccattcactt gcaaggcttt tgtttttgat aaagcaagaa aacaatgcct ctggttcccc      300
425 ttcaatagca tgtcaagtgg agtgaaaaaa gaatttggcc atgaatttga cctctatgaa      360
427 aacaaagact acattagaaa ctgcatcatt ggtaaaggac gcagctacaa gggaacagta      420
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431 agctatcggg gtaaagacct acaggaaaac tactgtcgaa atcctcgagg ggaagaaggg      540
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435 tgttcagaag ttgaatgcat gacctgcaat ggggagagtt atcgaggtct catggatcat      660
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451 tttaccactg atccaaacat ccgagttggc tactgctccc aaattccaaa ctgtgatatg     1140
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**RAW SEQUENCE LISTING ERROR SUMMARY**    DATE: 12/05/2006
PATENT APPLICATION:  **US/10/582,973A**    TIME: 09:48:42

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<u>**Invalid <213> Response:**</u>
**Use of "Artificial" only as "<213> Organism" response is incomplete,**
**per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.**

Seq#:4,5,6,7,8

**VERIFICATION SUMMARY**          DATE: 12/05/2006
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L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date